

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/582,370
Source: IFWP
Date Processed by STIC: 8/7/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>**> , **EFS Submission User Manual - ePAVE**)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: <u>10/587,370</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 ____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 ____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 ____ Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.	
4 ____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 ____ Variable Length	Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 ____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 ____ Skipped Sequences (OLD RULES)	Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 ____ Skipped Sequences (NEW RULES)	Sequence(s) ____ missing. If intentional , please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 ____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa , and which residue n or Xaa represents.	
10 ____ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)	
11 ____ Use of <220>	Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules	
12 ____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 ____ Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>	



IFWP

RAW SEQUENCE LISTING

DATE: 08/07/2006

PATENT APPLICATION: US/10/587,370

TIME: 09:35:48

Input Set : A:\11669.0237USWO-Sequence Listing.txt

Output Set: N:\CRF4\08072006\J587370.raw

3 <110> APPLICANT: Genentech, Inc.
 5 <120> TITLE OF INVENTION: BCMA POLYPEPTIDES AND USES THEREOF
 7 <130> FILE REFERENCE: 11669.0237USWO
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/587,370
 10 <141> CURRENT FILING DATE: 2006-07-26
 12 <150> PRIOR APPLICATION NUMBER: PCT/US2004/025247
 13 <151> PRIOR FILING DATE: 2004-08-04
 15 <150> PRIOR APPLICATION NUMBER: US 60/540,271
 16 <151> PRIOR FILING DATE: 2004-01-29
 18 <160> NUMBER OF SEQ ID NOS: 33
 20 <170> SOFTWARE: PatentIn version 3.3
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 34
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Artificial Sequence
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: Formula I
 31 <220> FEATURE:
 32 <221> NAME/KEY: MISC_FEATURE
 33 <222> LOCATION: (2)..(5)
 34 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 36 <220> FEATURE:
 37 <221> NAME/KEY: MISC_FEATURE
 38 <222> LOCATION: (7)..(7)
 39 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 41 <220> FEATURE:
 42 <221> NAME/KEY: MISC_FEATURE
 43 <222> LOCATION: (9)..(9)
 44 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 46 <220> FEATURE:
 47 <221> NAME/KEY: MISC_FEATURE
 48 <222> LOCATION: (11)..(11)
 49 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
 51 <220> FEATURE:
 52 <221> NAME/KEY: MISC_FEATURE
 53 <222> LOCATION: (12)..(13)
 54 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 56 <220> FEATURE:
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 58 <222> LOCATION: (16)..(16)
 59 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 61 <220> FEATURE:
 62 <221> NAME/KEY: MISC_FEATURE

ppr 1-6

Does Not Comply
Corrected Diskette Needed

insufficient. Give
source of genetic
material
(see item 11
on Enon
summary
sheet)

RAW SEQUENCE LISTING

DATE: 08/07/2006

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TIME: 09:35:48

Input Set : A:\11669.0237USWO-Sequence Listing.txt

Output Set: N:\CRF4\08072006\J587370.raw

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63 <222> LOCATION: (18)..(20)
64 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
66 <220> FEATURE:
67 <221> NAME/KEY: MISC_FEATURE
68 <222> LOCATION: (22)..(29)
69 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
71 <220> FEATURE:
72 <221> NAME/KEY: MISC_FEATURE
73 <222> LOCATION: (31)..(33)
74 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
76 <400> SEQUENCE: 1
W--> 78 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
      79 1          5          10          15
      82 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
      83          20          25          30
      86 Xaa Cys
90 <210> SEQ ID NO: 2
91 <211> LENGTH: 40
92 <212> TYPE: PRT
93 <213> ORGANISM: Artificial Sequence
95 <220> FEATURE:
96 <223> OTHER INFORMATION: Formula I
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105 <221> NAME/KEY: MISC_FEATURE
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107 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
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110 <221> NAME/KEY: MISC_FEATURE
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112 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
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115 <221> NAME/KEY: MISC_FEATURE
116 <222> LOCATION: (11)..(11)
117 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
119 <220> FEATURE:
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125 <221> NAME/KEY: MISC_FEATURE
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127 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
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130 <221> NAME/KEY: MISC_FEATURE
131 <222> LOCATION: (18)..(20)

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TIME: 09:35:48

Input Set : A:\11669.0237USWO-Sequence Listing.txt

Output Set: N:\CRF4\08072006\J587370.raw

132 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 134 <220> FEATURE:
 135 <221> NAME/KEY: MISC_FEATURE
 136 <222> LOCATION: (22)..(29)
 137 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 139 <220> FEATURE:
 140 <221> NAME/KEY: MISC_FEATURE
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 142 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
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 W--> 146 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
 147 1 5 10 15
 150 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 151 20 25 30
 154 Xaa Cys Asn Ser Val Lys Gly Thr
 155 35 40
 158 <210> SEQ ID NO: 3
 159 <211> LENGTH: 34
 160 <212> TYPE: PRT
 161 <213> ORGANISM: Artificial Sequence
 163 <220> FEATURE:
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 167 <220> FEATURE:
 168 <221> NAME/KEY: MISC_FEATURE
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 172 <220> FEATURE:
 173 <221> NAME/KEY: MISC_FEATURE
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 175 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
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 178 <221> NAME/KEY: MISC_FEATURE
 179 <222> LOCATION: (9)..(9)
 180 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 182 <220> FEATURE:
 183 <221> NAME/KEY: MISC_FEATURE
 184 <222> LOCATION: (11)..(11)
 185 <223> OTHER INFORMATION: Xaa is Leu, Val or Ile
 187 <220> FEATURE:
 188 <221> NAME/KEY: MISC_FEATURE
 189 <222> LOCATION: (12)..(13)
 190 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 192 <220> FEATURE:
 193 <221> NAME/KEY: MISC_FEATURE
 194 <222> LOCATION: (16)..(16)
 195 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 197 <220> FEATURE:
 198 <221> NAME/KEY: MISC_FEATURE
 199 <222> LOCATION: (18)..(20)

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Input Set : A:\11669.0237USWO-Sequence Listing.txt

Output Set: N:\CRF4\08072006\J587370.raw

200 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 202 <220> FEATURE:
 203 <221> NAME/KEY: MISC_FEATURE
 204 <222> LOCATION: (22)..(29)
 205 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 207 <220> FEATURE:
 208 <221> NAME/KEY: MISC_FEATURE
 209 <222> LOCATION: (31)..(33)
 210 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 212 <400> SEQUENCE: 3
 W--> 214 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
 215 1 5 10 15
 218 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 219 20 25 30
 222 Xaa Cys
 226 <210> SEQ ID NO: 4
 227 <211> LENGTH: 34
 228 <212> TYPE: PRT
 229 <213> ORGANISM: Artificial Sequence
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 232 <223> OTHER INFORMATION: Formula I
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 236 <221> NAME/KEY: MISC_FEATURE
 237 <222> LOCATION: (2)..(5)
 238 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 240 <220> FEATURE:
 241 <221> NAME/KEY: MISC_FEATURE
 242 <222> LOCATION: (7)..(7)
 243 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 245 <220> FEATURE:
 246 <221> NAME/KEY: MISC_FEATURE
 247 <222> LOCATION: (9)..(9)
 248 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 250 <220> FEATURE:
 251 <221> NAME/KEY: MISC_FEATURE
 252 <222> LOCATION: (11)..(11)
 253 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
 255 <220> FEATURE:
 256 <221> NAME/KEY: MISC_FEATURE
 257 <222> LOCATION: (12)..(13)
 258 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 260 <220> FEATURE:
 261 <221> NAME/KEY: MISC_FEATURE
 262 <222> LOCATION: (16)..(16)
 263 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 265 <220> FEATURE:
 266 <221> NAME/KEY: MISC_FEATURE
 267 <222> LOCATION: (18)..(18)
 268 <223> OTHER INFORMATION: Xaa is selected from the group consisting of Gln, Asp and

Ala

RAW SEQUENCE LISTING

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TIME: 09:35:48

Input Set : A:\11669.0237USWO-Sequence Listing.txt

Output Set: N:\CRF4\08072006\J587370.raw

270 <220> FEATURE:
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 272 <222> LOCATION: (19)..(20)
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 277 <222> LOCATION: (22)..(29)
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 282 <222> LOCATION: (31)..(33)
 283 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 285 <400> SEQUENCE: 4
 W--> 287 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
 288 1 5 10 15
 291 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 292 20 25 30
 295 Xaa Cys
 299 <210> SEQ ID NO: 5
 300 <211> LENGTH: 34
 301 <212> TYPE: PRT
 302 <213> ORGANISM: Artificial Sequence
 304 <220> FEATURE:
 305 <223> OTHER INFORMATION: Formula I *please correct this error in subsequent sequences.*
 308 <220> FEATURE:
 309 <221> NAME/KEY: MISC_FEATURE
 310 <222> LOCATION: (2)..(5)
 311 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
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 316 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 318 <220> FEATURE:
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 323 <220> FEATURE:
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 325 <222> LOCATION: (11)..(11)
 326 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
 328 <220> FEATURE:
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 336 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 338 <220> FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/587,370

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 TIME: 09:35:49

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 Output Set: N:\CRF4\08072006\J587370.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. ~~2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29~~
 Seq#:1; Xaa Pos. ~~31,32,33~~
 Seq#:2; Xaa Pos. ~~2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29~~
 Seq#:2; Xaa Pos. ~~31,32,33~~
 Seq#:3; Xaa Pos. ~~2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29~~
 Seq#:3; Xaa Pos. ~~31,32,33~~
 Seq#:4; Xaa Pos. ~~2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29~~
 Seq#:4; Xaa Pos. ~~31,32,33~~
 Seq#:5; Xaa Pos. ~~2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32~~
 Seq#:5; Xaa Pos. 33
 Seq#:10; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
 Seq#:10; Xaa Pos. 28,29,31,32,33
 Seq#:11; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
 Seq#:11; Xaa Pos. 28,29,31,32,33
 Seq#:12; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
 Seq#:12; Xaa Pos. 28,29,31,32,33
 Seq#:19; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
 Seq#:19; Xaa Pos. 28,29,31,32,33
 Seq#:28; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,19,22,23,24,25,26,27,28,29,31
 Seq#:28; Xaa Pos. 32,33
 Seq#:33; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
 Seq#:33; Xaa Pos. 28,29,31,32,33

VERIFICATION SUMMARY

DATE: 08/07/2006

PATENT APPLICATION: US/10/587,370

TIME: 09:35:49

Input Set : A:\11669.0237USWO-Sequence Listing.txt

Output Set: N:\CRF4\08072006\J587370.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
M:341 Repeated in SeqNo=1
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
M:341 Repeated in SeqNo=2
L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
M:341 Repeated in SeqNo=3
L:287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
M:341 Repeated in SeqNo=4
L:355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
M:341 Repeated in SeqNo=5
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
M:341 Repeated in SeqNo=10
L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
M:341 Repeated in SeqNo=11
L:758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
M:341 Repeated in SeqNo=12
L:995 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
M:341 Repeated in SeqNo=19
L:1485 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
M:341 Repeated in SeqNo=28
L:1734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
M:341 Repeated in SeqNo=33